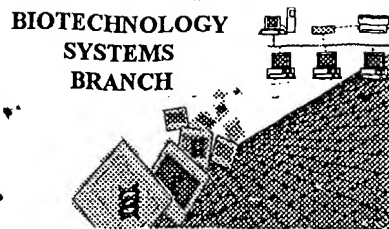# RAW SEQUENCE LISTING
ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: _10/615 479_

Source: _I fuo_

Date Processed by STIC: _12/10/03_

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:
1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 703-308-4212; FAX: 703-308-4221
Effective 12/13/03: TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE <u>CHECKER VERSION 4.1 PROGRAM</u>, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:
<u>http://www.uspto.gov/web/offices/pac/checker/chkr41note.htm</u>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom. Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<u><http://www.uspto.gov/ebc/efs/downloads/documents.htm></u> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry directly to (EFFECTIVE 12/01/03):
   U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 10/08/03

Raw Sequence Listing Error Summary

ERROR DETECTED   **SUGGESTED CORRECTION**   SERIAL NUMBER: 10/615,479

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

1 ____ Wrapped Nucleics   The number/text at the end of each line "wrapped" down to the next line. This may occur if your file
  Wrapped Aminos   was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will
                            prevent "wrapping."

2 ____ Invalid Line Length The rules require that a line not exceed 72 characters in length. This includes white spaces.

3 ____ Misaligned Amino   The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers;
  Numbering   use space characters, instead.

4 ____ Non-ASCII   The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please
                            ensure your subsequent submission is saved in ASCII text.

5 ____ Variable Length   Sequence(s) ____ contain n's or Xaa's representing more than one residue. Per Sequence Rules,
                            each n or Xaa can only represent a single residue. Please present the maximum number of each
                            residue having variable length and indicate in the <220>-<223> section that some may be missing.

6 ____ PatentIn 2.0   A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid
  "bug"   sequences(s) _____. Normally, PatentIn would automatically generate this section from the
                            previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to
                            the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for
                            Artificial or Unknown sequences.

7 ____ Skipped Sequences   Sequence(s) ____ missing. If intentional, please insert the following lines for each skipped sequence:
  (OLD RULES)   (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
                            (i)   SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
                            (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
                            This sequence is intentionally skipped

                            Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.

8 ____ Skipped Sequences   Sequence(s) ____ missing. If intentional, please insert the following lines for each skipped sequence.
  (NEW RULES)   <210> sequence id number
                            <400> sequence id number
                            000

9 ____ Use of n's or Xaa's Use of n's and/or Xaa's have been detected in the Sequence Listing.
  (NEW RULES)   Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
                            In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.

10 ____ Invalid <213>   Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or
  Response   scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or
                            is Artificial Sequence

11 ____ Use of <220>   Sequence(s) _____ missing the <220> "Feature" and associated numeric identifiers and responses.
                            Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or
                            "Unknown." Please explain source of genetic material in <220> to <223> section.
                            (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32)   (Sec. 1.823 of Sequence Rules)

12 ____ PatentIn 2.0   Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file,
  "bug"   resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence
                            listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.

13 ____ Misuse of n/Xaa   "n" can only represent a single <u>nucleotide</u>; "Xaa" can only represent a single <u>amino acid</u>

AMC – Biotechnology Systems Branch – 09/09/2003

IFWO

RAW SEQUENCE LISTING
PATENT APPLICATION: **US/10/615,479**

DATE: 12/10/2003
TIME: 17:33:46

Input Set : **A:\PTO.YF.txt**
Output Set: **N:\CRF4\12102003\J615479.raw**

```
 3 <110> APPLICANT: Larson Mr., Richard S.
 5 <120> TITLE OF INVENTION: Peptide Inhibitors of LFA-1/ICAM-1 Interaction
 7 <130> FILE REFERENCE: N12-012.CIP1
 9 <140> CURRENT APPLICATION NUMBER: 10/615,479
10 <141> CURRENT FILING DATE: 2003-07-08
12 <160> NUMBER OF SEQ ID NOS: 33
14 <170> SOFTWARE: Patent In Ver. 4.1
```

ERRORED SEQUENCES

Does Not Comply
Corrected Diskette Needed
p 1-5

```
      16 <210> SEQ ID NO: 1
      17 <211> LENGTH: 9
      18 <212> TYPE: PRT
      19 <213> ORGANISM: Artificial Sequence
      21 <220> FEATURE:
      22 <223> OTHER INFORMATION: Description of Artificial Sequence:Synthetic
      24 <400> SEQUENCE: 1
      25 Cys Leu Leu Arg Met Arg Ser Arg Cys                                9
E--> 26 1                    5              5
      28 <210> SEQ ID NO: 2
      29 <211> LENGTH: 9
      30 <212> TYPE: PRT
      31 <213> ORGANISM: Artificial Sequence
      33 <220> FEATURE:
      34 <223> OTHER INFORMATION: Description of Artificial Sequence:Synthetic
      36 <400> SEQUENCE: 2
      37 Cys Leu Leu Arg Met Arg Leu Ile Cys
E--> 38 1                    5              5
      40 <210> SEQ ID NO: 3
      41 <211> LENGTH: 9
      42 <212> TYPE: PRT
      43 <213> ORGANISM: Artificial Sequence
      45 <220> FEATURE:
      46 <223> OTHER INFORMATION: Description of Artificial Sequence:Synthetic
      49 <400> SEQUENCE: 3
      50 Cys Leu Leu Arg Met Arg Ser Ile Cys                                9
E--> 51 1    1          5                   5
      52 <210> SEQ ID NO: 4
      53 <211> LENGTH: 9
      54 <212> TYPE: PRT
      55 <213> ORGANISM: Artificial Sequence
      57 <220> FEATURE:
```

misAligned Amino Acid numbers. See item 3 on Error Summary Sheet.

RAW SEQUENCE LISTING   DATE: 12/10/2003
PATENT APPLICATION: **US/10/615,479**   TIME: 17:33:46

Input Set : **A:\PTO.YF.txt**
Output Set: **N:\CRF4\12102003\J615479.raw**

```
 58 <223> OTHER INFORMATION: Description of Artificial Sequence:Synthetic
 60 <400> SEQUENCE: 4
 61 Cys Leu Leu Arg Met Arg Leu Arg Cys
E--> 62 1                                                           9
 64 <210> SEQ ID NO: 5
 65 <211> LENGTH: 9
 66 <212> TYPE: PRT
 67 <213> ORGANISM: Artificial Sequence
 69 <220> FEATURE:
 70 <223> OTHER INFORMATION: Description of Artificial Sequence:Synthetic
 72 <400> SEQUENCE: 5
 73 Cys Leu Leu Arg Pro Arg Ser Ile Cys
E--> 74 1                                                           9
 76 <210> SEQ ID NO: 6
 77 <211> LENGTH: 9
 78 <212> TYPE: PRT
 79 <213> ORGANISM: Artificial Sequence
 81 <220> FEATURE:
 82 <223> OTHER INFORMATION: Description of Artificial Sequence:Synthetic
 84 <400> SEQUENCE: 6
 85 Cys Leu Leu Arg Pro Arg Ser Arg Cys                            9
E--> 86 1
 88 <210> SEQ ID NO: 7
 89 <211> LENGTH: 9
 90 <212> TYPE: PRT
 92 <213> ORGANISM: Artificial Sequence
 94 <220> FEATURE:
 95 <223> OTHER INFORMATION: Description of Artificial Sequence:Synthetic
 97 <400> SEQUENCE: 7
 98 Cys Leu Leu Arg Pro Arg Leu Ile Cys
E--> 99 1                                                           9
101 <210> SEQ ID NO: 8
102 <211> LENGTH: 9
103 <212> TYPE: PRT
104 <213> ORGANISM: Artificial Sequence
106 <220> FEATURE:
107 <223> OTHER INFORMATION: Description of Artificial Sequence:Synthetic
109 <400> SEQUENCE: 8
110 Cys Leu Leu Arg Pro Arg Leu Arg Cys
E--> 111 1                                                          9
113 <210> SEQ ID NO: 9
114 <211> LENGTH: 9
115 <212> TYPE: PRT
116 <213> ORGANISM: Artificial Sequence
118 <220> FEATURE:
119 <223> OTHER INFORMATION: Description of Artificial Sequence:Synthetic
121 <400> SEQUENCE: 9
122 Cys Leu Leu Arg Met Asn Ser Ile Cys                            9
E--> 123 1
```

Handwritten annotations: "Same error" (×6), with "5", "5", and "9" markings next to lines 62, 74, 86, 99, 111, 123.

```
        125 <210> SEQ ID NO: 10
        126 <211> LENGTH: 9
        127 <212> TYPE: PRT
        128 <213> ORGANISM: Artificial Sequence
        130 <220> FEATURE:
        131 <223> OTHER INFORMATION: Description of Artificial Sequence:Synthetic
        133 <400> SEQUENCE: 10
        134 Cys Leu Leu Arg Met Asn Ser Arg Cys                        9
E-->    135 1   .                        5
        137 <210> SEQ ID NO: 11
        138 <211> LENGTH: 9
        139 <212> TYPE: PRT
        140 <213> ORGANISM: Artificial Sequence
        142 <220> FEATURE:
        143 <223> OTHER INFORMATION: Description of Artificial Sequence:Synthetic
        145 <400> SEQUENCE: 11
        146 Cys Leu Leu Arg Met Asn Leu Ile Cys                        9
E-->    147 1                            5
        149 <210> SEQ ID NO: 12
        150 <211> LENGTH: 9
        151 <212> TYPE: PRT
        152 <213> ORGANISM: Artificial Sequence
        154 <220> FEATURE:
        155 <223> OTHER INFORMATION: Description of Artificial Sequence:Synthetic
        157 <400> SEQUENCE: 12
        158 Cys Leu Leu Arg Met Asn Leu Arg Cys                        9
E-->    159 1                            5
        161 <210> SEQ ID NO: 13
        162 <211> LENGTH: 9
        163 <212> TYPE: PRT
        164 <213> ORGANISM: Artificial Sequence
        166 <220> FEATURE:
        167 <223> OTHER INFORMATION: Description of Artificial Sequence:Synthetic
        169 <400> SEQUENCE: 13
        170 Cys Leu Leu Arg Pro Asn Ser Ile Cys                        9
E-->    171 1                            5
        173 <210> SEQ ID NO: 14
        174 <211> LENGTH: 9
        175 <212> TYPE: PRT
        176 <213> ORGANISM: Artificial Sequence
        178 <220> FEATURE:
        179 <223> OTHER INFORMATION: Description of Artificial Sequence:Synthetic
        181 <400> SEQUENCE: 14
        182 Cys Leu Leu Arg Pro Asn Ser Arg Cys                        9
E-->    183 1                            5
        185 <210> SEQ ID NO: 15
        186 <211> LENGTH: 9
        187 <212> TYPE: PRT
        188 <213> ORGANISM: Artificial Sequence
```

same error (handwritten annotation, with 9 and 5 circled)

same error (handwritten annotation)

same error (handwritten annotation)

same error (handwritten annotation)

same error (handwritten annotation)

RAW SEQUENCE LISTING   DATE: 12/10/2003
PATENT APPLICATION: US/10/615,479   TIME: 17:33:46

Input Set : **A:\PTO.YF.txt**
Output Set: **N:\CRF4\12102003\J615479.raw**

```
        190 <220> FEATURE:
        191 <223> OTHER INFORMATION: Description of Artificial Sequence:Synthetic.
        193 <400> SEQUENCE: 15
        194 Cys Leu Leu Arg Pro Asn Leu Ile Cys                            9
E-->    195 1                      5

        197 <210> SEQ ID NO: 16
        198 <211> LENGTH: 9
        199 <212> TYPE: PRT
        200 <213> ORGANISM: Artificial Sequence
        202 <220> FEATURE:
        203 <223> OTHER INFORMATION: Description of Artificial Sequence:Synthetic
        205 <400> SEQUENCE: 16
        206 Cys Leu Leu Arg Pro Asn Leu Arg Cys                            9
E-->    207 1                      5

        209 <210> SEQ ID NO: 17
        210 <211> LENGTH: 9
        211 <212> TYPE: PRT
        212 <213> ORGANISM: Artificial Sequence
        214 <220> FEATURE:
        215 <223> OTHER INFORMATION: Description of Artificial Sequence:Synthetic
        217 <400> SEQUENCE: 17
        218 Cys Met Leu Arg Met Arg Ser Ile Cys                            9
E-->    219 1                      5

        221 <210> SEQ ID NO: 18
        222 <211> LENGTH: 9
        223 <212> TYPE: PRT
        224 <213> ORGANISM: Artificial Sequence
        227 <220> FEATURE:
        228 <223> OTHER INFORMATION: Description of Artificial Sequence:Synthetic
        230 <400> SEQUENCE: 18
        231 Cys Met Leu Arg Met Arg Ser Arg Cys                            9
E-->    232 1                      5

        234 <210> SEQ ID NO: 19
        235 <211> LENGTH: 9
        236 <212> TYPE: PRT
        237 <213> ORGANISM: Artificial Sequence
        239 <220> FEATURE:
        240 <223> OTHER INFORMATION: Description of Artificial Sequence:Synthetic
        242 <400> SEQUENCE: 19
        243 Cys Met Leu Arg Met Arg Leu Ile Cys                            9
E-->    244 1                      5
        246 <210> SEQ ID NO: 20
        247 <211> LENGTH: 9
        248 <212> TYPE: PRT
        249 <213> ORGANISM: Artificial Sequence
        251 <220> FEATURE:
        252 <223> OTHER INFORMATION: Description of Artificial Sequence:Synthetic
        254 <400> SEQUENCE: 20
        255 Cys Met Leu Arg Met Arg Leu Arg Cys                            9
```

[handwritten annotations: "SAme error" appears five times alongside lines 195, 207, 219, 232, and 244]

RAW SEQUENCE LISTING   DATE: 12/10/2003
PATENT APPLICATION: US/10/615,479   TIME: 17:33:46

Input Set : A:\PTO.YF.txt
Output Set: N:\CRF4\12102003\J615479.raw

E--> 256 1

VERIFICATION SUMMARY   DATE: 12/10/2003
PATENT APPLICATION: **US/10/615,479**   TIME: 17:33:47

Input Set : **A:\PTO.YF.txt**
Output Set: **N:\CRF4\12102003\J615479.raw**

```
L:26  M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:1
L:26  M:301 E: (44) No Sequence Data was Shown, SEQ ID:1
L:26  M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:1
L:38  M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:2
L:38  M:301 E: (44) No Sequence Data was Shown, SEQ ID:2
L:38  M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:2
L:51  M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:3
L:51  M:301 E: (44) No Sequence Data was Shown, SEQ ID:3
L:51  M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:3
L:62  M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:4
L:62  M:301 E: (44) No Sequence Data was Shown, SEQ ID:4
L:62  M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:4
L:74  M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:5
L:74  M:301 E: (44) No Sequence Data was Shown, SEQ ID:5
L:74  M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:5
L:86  M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:6
L:86  M:301 E: (44) No Sequence Data was Shown, SEQ ID:6
L:86  M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:6
L:99  M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7
L:99  M:301 E: (44) No Sequence Data was Shown, SEQ ID:7
L:99  M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:7
L:111 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:8
L:111 M:301 E: (44) No Sequence Data was Shown, SEQ ID:8
L:111 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:8
L:123 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:9
L:123 M:301 E: (44) No Sequence Data was Shown, SEQ ID:9
L:123 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:9
L:135 M:301 E: (44) No Sequence Data was Shown, SEQ ID:10
L:135 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:10
L:147 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:11
L:147 M:301 E: (44) No Sequence Data was Shown, SEQ ID:11
L:147 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:11
L:159 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:12
L:159 M:301 E: (44) No Sequence Data was Shown, SEQ ID:12
L:159 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:12
L:171 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:13
L:171 M:301 E: (44) No Sequence Data was Shown, SEQ ID:13
L:171 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:13
L:183 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:14
L:183 M:301 E: (44) No Sequence Data was Shown, SEQ ID:14
L:183 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:14
L:195 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:15
L:195 M:301 E: (44) No Sequence Data was Shown, SEQ ID:15
L:195 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:15
L:207 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:16
L:207 M:301 E: (44) No Sequence Data was Shown, SEQ ID:16
L:207 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:16
L:219 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:17
```

VERIFICATION SUMMARY   DATE: 12/10/2003
PATENT APPLICATION: **US/10/615,479**   TIME: 17:33:47

Input Set : **A:\PTO.YF.txt**
Output Set: **N:\CRF4\12102003\J615479.raw**

```
L:219 M:301 E: (44) No Sequence Data was Shown, SEQ ID:17
L:219 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:17
L:232 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:18
L:232 M:301 E: (44) No Sequence Data was Shown, SEQ ID:18
L:232 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:18
L:244 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:19
L:244 M:301 E: (44) No Sequence Data was Shown, SEQ ID:19
L:244 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:19
L:256 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:20
L:256 M:301 E: (44) No Sequence Data was Shown, SEQ ID:20
L:256 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:20
L:268 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:21
L:268 M:301 E: (44) No Sequence Data was Shown, SEQ ID:21
L:268 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:21
L:280 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:22
L:280 M:301 E: (44) No Sequence Data was Shown, SEQ ID:22
L:280 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:22
L:292 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:23
L:292 M:301 E: (44) No Sequence Data was Shown, SEQ ID:23
L:292 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:23
L:304 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:24
L:304 M:301 E: (44) No Sequence Data was Shown, SEQ ID:24
L:304 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:24
L:316 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:25
L:316 M:301 E: (44) No Sequence Data was Shown, SEQ ID:25
L:316 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:25
L:328 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:26
L:328 M:301 E: (44) No Sequence Data was Shown, SEQ ID:26
L:328 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:26
L:340 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:27
L:340 M:301 E: (44) No Sequence Data was Shown, SEQ ID:27
L:340 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:27
L:352 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:28
L:352 M:301 E: (44) No Sequence Data was Shown, SEQ ID:28
L:352 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:28
L:364 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:29
L:364 M:301 E: (44) No Sequence Data was Shown, SEQ ID:29
L:364 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:29
L:376 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:30
L:376 M:301 E: (44) No Sequence Data was Shown, SEQ ID:30
L:376 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:30
L:388 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:31
L:388 M:301 E: (44) No Sequence Data was Shown, SEQ ID:31
L:388 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:31
L:400 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:32
L:400 M:301 E: (44) No Sequence Data was Shown, SEQ ID:32
L:400 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:32
L:414 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:33
L:414 M:301 E: (44) No Sequence Data was Shown, SEQ ID:33
```

VERIFICATION SUMMARY　　　　　　　　DATE: 12/10/2003
PATENT APPLICATION:　**US/10/615,479**　　TIME: 17:33:47

Input Set : **A:\PTO.YF.txt**
Output Set: **N:\CRF4\12102003\J615479.raw**

L:414 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:0 SEQ:33